

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	/0/628,296
Source:	OHE
Date Processed by STIC:	8.15/2003

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<a href="http://www.uspto.gov/ebc/efs/downloads/documents.htm">http://www.uspto.gov/ebc/efs/downloads/documents.htm</a>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450

Additional Actions

- 3. Hand Carry directly to:
  - U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
  - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003



OIPE

RAW SEQUENCE LISTING DATE: 08/05/2003 PATENT APPLICATION: US/10/628,296 TIME: 10:08:31

Input Set : A:\P06331US01.ST25.txt

Output Set: N:\CRF4\08052003\J628296.raw

```
5 <110> APPLICANT: Li, Lian-Chao
             Cosgrove, Daniel J.
    11 <120> TITLE OF INVENTION: PLANT CELL WALL LOOSENING ACTIVITY OF GROUP 2/3 ALLERGENS OF
GRASS POLLEN
     15 <130> FILE REFERENCE: P06331US01
C--> 19 <140> CURRENT APPLICATION NUMBER: US/10/628,296
     20 <141> CURRENT FILING DATE: 2003-07-28
     24 <150> PRIOR APPLICATION NUMBER: US 60,399,688
     26 <151> PRIOR FILING DATE: 2002-07-29
     30 <160> NUMBER OF SEQ ID NOS: 18
                                                             Does Not Comply
     34 <170> SOFTWARE: PatentIn version 3.1
     38 <210> SEQ ID NO: 1
                                                         Corrected Diskette Needed
     40 <211> LENGTH: 291
                                                        pg 1-3,6
    42 <212> TYPE: DNA
    44 <213> ORGANISM: Lolium perenne
    48 <400> SEQUENCE: 1
    49 acaaaagtcg atttaactgt ggagaagggt tctgacgcga agacgctggt gctgaacatc
                                                                               60
     51 aagtacacga ggccagggga caccctggcg gaggtggagc teeggcagca eggeteggag
    53 gagtgggaac ccatgacgaa gaagggcaac ctgtgggagg tgaagagcgc caagccgctc
                                                                             180
    55 accggcccaa tgaacttccg cttcctctcc aagggcggca tgaagaacgt cttcgacgag
                                                                             240
    57 gtcatcccca ccgccttcac ggtcggcaaa acctacaccc cagaatacaa t
                                                                             291
    60 <210> SEQ ID NO: 2
    62 <211> LENGTH: 97
    64 <212> TYPE: PRT
    66 <213> ORGANISM: Lolium perenne
    70 <400> SEQUENCE: 2
    72 Thr Lys Val Asp Leu Thr Val Glu Lys Gly Ser Asp Ala Lys Thr Leu
    73 1
                                            10
    76 Val Leu Asn Ile Lys Tyr Thr Arg Pro Gly Asp Thr Leu Ala Glu Val
    77
                                        25
    80 Glu Leu Arg Gln His Gly Ser Glu Glu Trp Glu Pro Met Thr Lys Lys
                                    40
    84 Gly Asn Leu Trp Glu Val Lys Ser Ala Lys Pro Leu Thr Gly Pro Met
    85
    88 Asn Phe Arg Phe Leu Ser Lys Gly Gly Met Lys Asn Val Phe Asp Glu
                            70
    92 Val Ile Pro Thr Ala Phe Thr Val Gly Lys Thr Tyr Thr Pro Glu Tyr
    93
    96 Asn
    100 <210> SEQ ID NO: 3
    102 <211> LENGTH: 291
    104 <212> TYPE: DNA
                                    sel p. 6 for enor explanation
    106 <213 ORGANISM: Artificial
```

60

120

180

240

291

PATENT APPLICATION: US/10/628,296

DATE: 08/05/2003
TIME: 10:08:31

Input Set: A:\P06331US01.ST25.txt
Output Set: N:\CRF4\08052003\J628296.raw

```
W--> 110 <223 OTHER INFORMATION:
W--> 110 < 400> 3
     111 acaaaagtcg atttaactgt ggagaagggt tctgacgcga agacgctggt gctgaacatc
     113 aagtacacga ggccagggga caccctggcg gaggtggagc tccggcagca cggctcggag
     115 gagtgggaac ccctgacgaa gaagggcaac ctgtgggagg tgaagagcgc caagccgctc
     117 accggcccaa tgaacttccg cttcctctcc aagggcggca tgaagaacgt cttcgacgag
     119 gtcatececa eegeetteae ggteggcaaa acetacaeee cagaatacaa t
     122 <210> SEQ ID NO: 4
     124 <211> LENGTH: 97
     126 <212> TYPE: PRT
128 <213> ORGANISM Artificial W--> 132 x220> FEATURE:
                                      see P.6
W--> 132 (<223) OTHER INFORMATION:
W--> 132 <400> 4
     134 Thr Lys Val Asp Leu Thr Val Glu Lys Gly Ser Asp Ala Lys Thr Leu
     135 1
     138 Val Leu Asn Ile Lys Tyr Thr Arg Pro Gly Asp Thr Leu Ala Glu Val
                  . 20
     142 Glu Leu Arg Gln His Gly Ser Glu Glu Trp Glu Pro Met Thr Lys Lys
                                      40
     146 Gly Asn Leu Trp Glu Val Lys Ser Ala Lys Pro Leu Thr Gly Pro Met
     150 Asn Phe Arg Phe Leu Ser Lys Gly Gly Met Lys Asn Val Phe Asp Glu
                             70
                                                  75
     154 Val Ile Ala Thr Ala Phe Thr Val Gly Lys Thr Tyr Thr Pro Glu Tyr
     155
     158 Asn
     162 <210> SEQ ID NO: 5
     164 <211> LENGTH: 20
     166 <212> TYPE: PRT
     168 <213> ORGANISM: Lolium perenne
     172 <400> SEQUENCE: 5
     174 Thr Lys Val Asp Leu Thr Val Glu Lys Gly Ser Asp Ala Lys Thr Leu
     175 1
                         5
                                              10
     178 Val Leu Asn Ile
     179
     182 <210> SEQ ID NO: 6
     184 <211> LENGTH: 52
     188 <213 > ORGANISM: Artificial needs explanation - sel p 6
192 <220 > FEATURE
     192 <220> FEATURE:
     194 <221> NAME/KEY: misc_feature
     196 <222> LOCATION: (3)..(3)
     198 <223> OTHER INFORMATION: N is unknown
     202 <220> FEATURE:
     204 <221> NAME/KEY: misc_feature
     206 <222> LOCATION: (15)..(15)
     208 <223> OTHER INFORMATION: N is unknown
     212 <220> FEATURE:
```

DATE: 08/05/2003

TIME: 10:08:31

```
Input Set : A:\P06331US01.ST25.txt
                         Output Set: N:\CRF4\08052003\J628296.raw
      214 <221> NAME/KEY: misc_feature
      216 <222> LOCATION: (29)..(29)
      218 <223> OTHER INFORMATION: N is unknown
      222 <220> FEATURE:
      224 <221> NAME/KEY: misc_feature
      226 <222> LOCATION: (36)..(36)
      228 <223> OTHER INFORMATION: N is unknown
      232 <220> FEATURE:
     234 <221> NAME/KEY: misc_feature
     236 <222> LOCATION: (43)..(43)
     238 <223> OTHER INFORMATION: N is unknown
     242 <400> SEQUENCE: 6
                                                                                               52
W--> 243 acnacgtaar aggtnacgtg ayctycttna cgtacnacgt gtnacgtgar ag
      246 <210> SEQ ID NO: 7
      248 <211> LENGTH: 49
     250 <212> TYPE: DNA
252 <213> ORGANISM Artificial
256 <220> FEATURE:
258 <221> NAME/KEY: misc_feature
260 <222> LOCATION: (24)..(24)
262 <223> OTHER INFORMATION: N is unknown
266 <220> FEATURE:
268 <221> NAME/KEY: misc_feature
270 <222> LOCATION: (31)..(31)
272 <223> OTHER INFORMATION: N is unknown
267 <223> OTHER INFORMATION: N is unknown
270 <225 LOCATION: (31)..(31)
     250 <212> TYPE: DNA
                                                                or of Unknown are equality
     272 <223> OTHER INFORMATION: N is unknown
     276 <220> FEATURE:
     278 <221> NAME/KEY: misc_feature
     280 <222> LOCATION: (43)...(43)
     282 <223> OTHER INFORMATION: N is unknown
     286 <400> SEQUENCE: 7
W--> 287 cyctyctara gttragtayc ttcnacgtgg nacgtgtrag tanacgtgt
      290 <210> SEQ ID NO: 8
      292 <211> LENGTH: 93
     294 <212> TYPE: PRT
     296 <213> ORGANISM: Zea mays
     300 <400> SEQUENCE: 8
     302 Thr Phe Gln Val Gly Lys Gly Ser Lys Pro Gly His Leu Val Leu Thr
     306 Pro Asn Ile Ala Thr Ile Ser Asp Val Glu Ile Lys Glu His Gly Gly
                        20
     310 Asp Asp Phe Ser Phe Thr Leu Lys Glu Gly Pro Ala Gly Thr Trp Thr
                   35
                                           40
     314 Leu Asp Thr Lys Ala Pro Leu Lys Tyr Pro Leu Cys Ile Arg Phe Ala
                                       55
                                                               60
     318 Thr Lys Ser Gly Gly Tyr Arg Ile Ala Asp Asp Val Ile Pro Ala Asp
     322 Phe Lys Ala Gly Thr Thr Tyr Lys Thr Thr Leu Ser Ile
     326 <210> SEQ ID NO: 9
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/628,296

RAW SEQUENCE LISTING DATE: 08/05/2003 PATENT APPLICATION: US/10/628,296 TIME: 10:08:31

Input Set : A:\P06331US01.ST25.txt
Output Set: N:\CRF4\08052003\J628296.raw

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328 <211> LENGTH: 96
330 <212> TYPE: PRT
332 <213> ORGANISM: Dactylis glomerata
336 <400> SEQUENCE: 9
338 Val Lys Val Thr Phe Lys Val Glu Lys Gly Ser Asp Pro Lys Lys Leu
342 Val Leu Asp Ile Lys Tyr Thr Arg Pro Gly Asp Thr Leu Ala Glu Val
346 Glu Leu Arg Gln His Gly Ser Glu Glu Trp Glu Pro Leu Thr Lys Lys
           35
                                40
350 Gly Asn Leu Trp Glu Val Lys Ser Ser Lys Pro Leu Thr Gly Pro Phe
                            55
354 Asn Phe Arg Phe Met Ser Lys Gly Gly Met Arg Asn Val Phe Asp Glu
                                            75
355 65
358 Val Ile Pro Thr Ala Phe Lys Ile Gly Thr Thr Tyr Thr Pro Glu Glu
359
362 <210> SEQ ID NO: 10
364 <211> LENGTH: 90
366 <212> TYPE: PRT
368 <213> ORGANISM: Oryza sativa
372 <400> SEQUENCE: 10
374 Met Glu Val Ala Lys Gly Ser Ser Ala Lys Ser Leu Glu Leu Val Thr
378 Asn Val Ala Ile Ser Lys Val Glu Val Lys Glu Lys Gly Gly Lys Asp
                                    25
                20
382 Trp Val Ala Leu Lys Glu Ser Ser Ser Asn Thr Trp Thr Leu Lys Ser
                                40
386 Glu Ser Pro Leu Lys Gly Pro Phe Ser Val Arg Phe Leu Val Lys Asn
                            55
390 Ser Gly Tyr Arg Val Val Asp Asp Ile Ile Pro Glu Ser Phe Thr Ala
394 Gly Ser Glu Tyr Lys Ser Gly Ile Gln Leu
395
                    85
398 <210> SEQ ID NO: 11
400 <211> LENGTH: 96
402 <212> TYPE: PRT
404 <213> ORGANISM: Lolium perenne
408 <400> SEQUENCE: 11
410 Ala Ala Pro Val Glu Phe Thr Val Glu Lys Gly Ser Asp Glu Lys Asn
414 Leu Ala Leu Ser Ile Lys Tyr Asn Lys Glu Gly Asp Ser Met Ala Glu
                20
                                    25
418 Val Glu Leu Lys Glu His Gly Ser Asn Glu Trp Leu Ala Leu Lys Lys
422 Asn Gly Asp Gly Val Trp Glu Ile Lys Ser Asp Lys Pro Leu Lys Gly
423
426 Pro Phe Asn Phe Arg Phe Val Ser Glu Lys Gly Met Arg Asn Val Phe
430 Asp Asp Val Val Pro Ala Asp Phe Lys Val Gly Thr Thr Tyr Lys Pro
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RAW SEQUENCE LISTING DATE: 08/05/2003 PATENT APPLICATION: US/10/628,296 TIME: 10:08:31

Input Set : A:\P06331US01.ST25.txt
Output Set: N:\CRF4\08052003\J628296.raw

431					85					90					95	
434	<210	)> SI	EQ II	ON C	: 12						•	•				
436	<211	l> LI	ENGTI	H: 90	5			•								
438	<212	2> TY	YPE:	PRT												
440	<213	3> OI	RGAN	ISM:	Sorg	ghum										
	<400				12					•						•
446	Gly	Thr	Thr	Leu	Thr	Ile	Glu	Val.	Gly	Lys	Asp	Ser	Thr	Ser	Thr	Lys
447	1				5				_	10	_				15	
450	Leu	Ser	Leu	Ile	Thr	Asn	Val	Ala	Ile	Ser	Glu	Val	Ser	Val	Lys	Pro
451				20					25					30		
454	Lys	Gly	Ala	Thr	Asp	Phe	Thr	Asp	Asp	Leu	Lys	Glu	Ser	Glu	Pro	Lys
455	•	-	35		~			40	_				45			
458	Thr	Phe	Thr	Leu	Asp	Ser	Lys	Glu	Pro	Ile	Glu	Gly	Pro	Ile	Ala	Phe
459		50			-		55					60				
462	Arg	Phe	Leu	Ala	Lys	Gly	Gly	Gly	Tyr	Arg	Val	Val	Asp	Asn	Ala	Ile
463					-	70		_	_	_	75		_			80
466	Pro	Ala	Asp	Phe	Lys	Ala	Gly	Ser	Va1	Tyr	Lys	Thr	Thr	Glu	Gln	Val
467			•		85		-			90	-				95	
470	O <210> SEQ ID NO: 13															
	<211															
	<212															
	<213				Horo	deum	vul	gare								
	<400				13											
	Ala					Lvs	Phe	Thr	Val	Gln	Lys	Gly	Ser	Asp	Ala	Lys
483				-	5	<b>-</b> .				10		-		-	15	-
	Lys	Leu	Val	Leu	Lys	Ile	Asp	Tyr	Thr	Arg	Ala	Gly	Asp	Thr	Leu	Ser
487	-			20	-		-	-	25	_		-	_	30		
490	Glu	Met	Glu	Leu	Arg	Gln	His	Gly	Ser	Glu	Glu	Trp	Glu	Pro	Phe	Thr
491			35		_			40				_	45			
494	Lys	Lys	Gly	Asp	Val	Trp	Glu	Leu	Ser	Ser	Ser	Lys	Pro	Leu	Val	Gly
495	-,	50	-	_		_	55					60				
498	Pro	Phe	Asn	Phe	Arg	Phe	Leu	Ser	Lys	Gly	Gly	Met	Lys	Asn	Val	Phe
499				•	_	70					75					80
502	Asp	Glu	Val	Phe	Ser	Thr	Asp	Phe	Lys	Ile	Gly	Lys	Thr	Tyr	Glu	Pro
503	_				85				_	90					95	
506	Val	Tyr	Asp	Ala												
507		_	-	100											•	
510	<210	)> SI	EQ II	ON C	: 14											
512	<211	l> Li	ENGT	H: 98	3											
514	<212	2> T	PE:	PRT												
516	<213	3> OI	RGAN.	ISM:	Trit	icur	n aes	stivu	ım							
	<400															
	Ala					Ser	Ile	Thr	Val	Glu	Lys	Gly	Ser	Asp	Ala	Lys
523					5					10		-		-	15	-
	His	Leu	Val	Leu		Ile	Lys	Tyr	Asp	Lys	Val	Gly	Asp	Ser	Met	Lys
527				20			-	-	25	-		-	-	30		-
	Glu	Val	Glu		Glu	Gln	Asn	Glu	Asp	Trp	Leu	Pro	Leu	Lys	Lys	Gly
531			35					40	_	-			45	-	-	-
	Tyr	Ser		Ala	Trp	Glu	Ile	Lys	Ser	Asp	Thr	Pro	Leu	Lys	Gly	Pro
	-	•	-		_			-		-				_	_	

RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 08/05/2003

PATENT APPLICATION: US/10/628,296

TIME: 10:08:32

Input Set : A:\P06331US01.ST25.txt

Output Set: N:\CRF4\08052003\J628296.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; N Pos. 3, 15, 2/9, 36, 4/8Seq#:7; N Pos. 24,31,43

Seq#:17; Xaa Pos. 27

## Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 11

## Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,4,6,7

Use of <220> Feature(NEW RULES): INO Applaration
Sequence(s) are missing the Sequence(s)\_are missing the  $\langle 220 \rangle$  Feature and associated headings. Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or"Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104,pp.29631-32) (Sec.1.823 of new Rules)

Seq#:3,4,6

file://C:\CRF4\Outhold\VsrJ628296.htm

VERIFICATION SUMMARY

DATE: 08/05/2003 PATENT APPLICATION: US/10/628,296 TIME: 10:08:32

Input Set : A:\P06331US01.ST25.txt

Output Set: N:\CRF4\08052003\J628296.raw

- L:19 M:270 C: Current Application Number differs, Wrong Format
- L:110 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:3, <213>
- ORGANISM: Artificial
- L:110 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:3, <213>
- ORGANISM: Artificial
- L:110 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:110
- L:132 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:4, <213>
- ORGANISM: Artificial
- L:132 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:4, <213> ORGANISM: Artificial
- L:132 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:132
- L:243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
- L:287 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0 L:652 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:16